

**A**

**pT-1 serine analogues abolish Plk1 PBD : peptide binding in solution**

Peptide name	Peptide sequence	$K_d$
PoloBoxide-optimal	MAGPMQ-S-pT-P-LNGAKK (SEQ ID NO: 3)	280 ± 27 nM
PoloBoxide-7A	MAGPMQ-A-pT-P-LNGAYKK (SEQ ID NO: 68)	N.D.B.
PoloBoxide-7G	MAGPMQ-G-pT-P-LNGAYKK (SEQ ID NO: 69)	N.D.B.
PoloBoxide-7C	MAGPMQ-C-pT-P-LNGAYKK (SEQ ID NO: 70)	N.D.B.
PoloBoxide-7T	MAGPMQ-T-pT-P-LNGAYKK (SEQ ID NO: 71)	N.D.B.

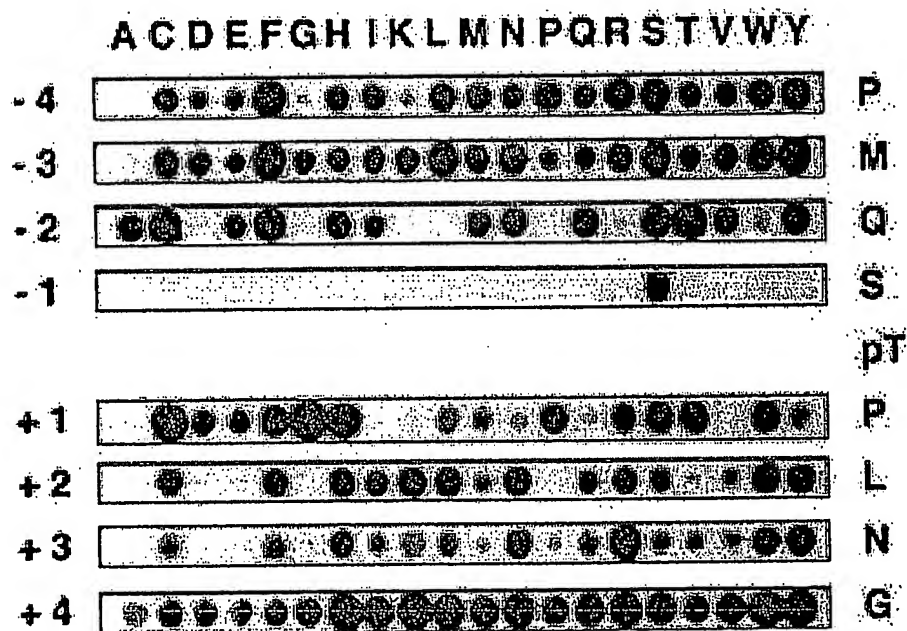
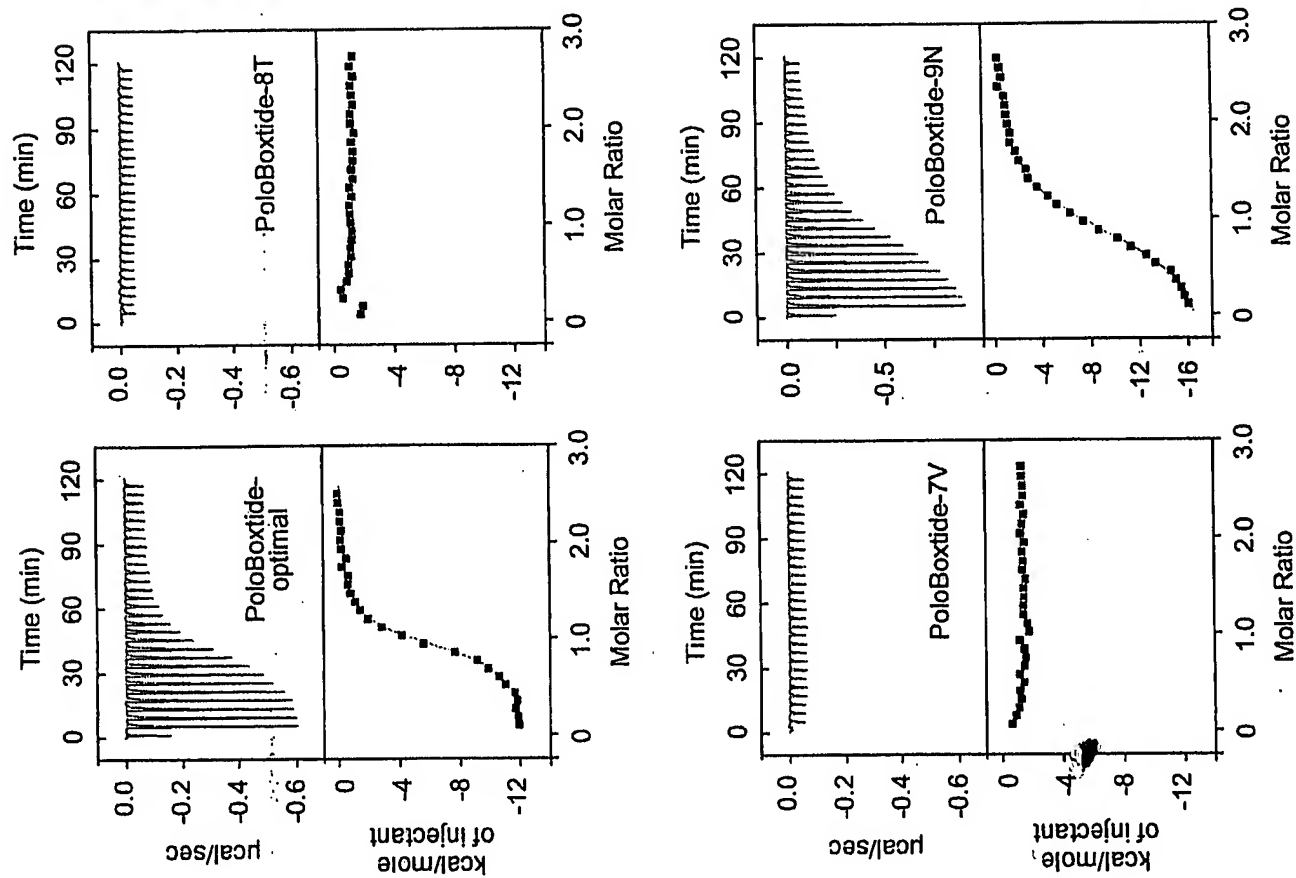
**B**

Figure 9



Peptide binding affinities for the Plk1 Polo Box Domain			
Peptide name	Peptide sequence	$K_d$	
PoloBoxide-optimal	MAGPMQ-S-pT-P-LNGAKK (SEQ ID NO: 3)	$280 \pm 27$ nM	
Effect of pT			
PoloBoxide-8T	MAGPMQ-S- <b>T</b> -P-LNGAKK (SEQ ID NO: 34)	N.D.B.	
PoloBoxide-8pS	MAGPMQ-S- <b>pS</b> -P-LNGAYKK (SEQ ID NO: 65)	2.1 $\mu$ M	
PoloBoxide-8pY	MAGPMQ-S- <b>pY</b> -P-LNGAYKK (SEQ ID NO: 66)	N.D.B.	
Effect of serine at pT-1 position			
PoloBoxide-7V	MAGPMQ- <b>V</b> -pT-P-LNGAKK (SEQ ID NO: 67)	N.D.B.	
PoloBoxide-7A	MAGPMQ- <b>A</b> -pT-P-LNGAYKK (SEQ ID NO: 68)	N.D.B.	
PoloBoxide-7G	MAGPMQ- <b>G</b> -pT-P-LNGAYKK (SEQ ID NO: 69)	N.D.B.	
PoloBoxide-7C	MAGPMQ- <b>C</b> -pT-P-LNGAYKK (SEQ ID NO: 70)	N.D.B.	
PoloBoxide-7T	MAGPMQ- <b>T</b> -pT-P-LNGAYKK (SEQ ID NO: 71)	N.D.B.	
Effect of proline at pT+1 position			
PoloBoxide-9N	MAGPMQ-S-pT- <b>N</b> -LNGAKK (SEQ ID NO: 72)	1.5 $\mu$ M	

Figure 4